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	500		505		510
Cys Gln Asp Val	Asp Gly Tyr Cys Tyr	Asn Gly Ile Cys Gln Thr			
	515		520		525
His Glu Gln Gln	Cys Val Thr Leu Trp	Gly Pro Gly Ala Lys Pro			
	530		535		540
Ala Pro Gly Ile	Cys Phe Glu Arg Val	Asn Ser Ala Gly Asp Pro			
	545		550		555
Tyr Gly Asn Cys	Gly Lys Val Ser Lys	Ser Ser Phe Ala Lys Cys			
	560		565		570
Glu Met Arg Asp	Ala Lys Cys Gly Lys	Ile Gln Cys Gln Gly Gly			
	575		580		585
Ala Ser Arg Pro	Val Ile Gly Thr Asn	Ala Val Ser Ile Glu Thr			
	590		595		600
Asn Ile Pro Leu	Gln Gln Gly Gly Arg	Ile Leu Cys Arg Gly Thr			
	605		610		615
His Val Tyr Leu	Gly Asp Asp Met Pro	Asp Pro Gly Leu Val Leu			
	620		625		630
Ala Gly Thr Lys	Cys Ala Asp Gly Lys	Ile Cys Leu Asn Arg Gln			
	635		640		645
Cys Gln Asn Ile	Ser Val Phe Gly Val	His Glu Cys Ala Met Gln			
	650		655		660
Cys His Gly Arg	Gly Val Cys Asn Asn	Arg Lys Asn Cys His Cys			
	665		670		675
Glu Ala His Trp	Ala Pro Pro Phe Cys	Asp Lys Phe Gly Phe Gly			
	680		685		690
Gly Ser Thr Asp	Ser Gly Pro Ile Arg	Gln Ala Glu Ala Arg Gln			
	695		700		705
Glu Ala Ala Glu	Ser Asn Arg Glu Arg	Gly Gln Gly Gln Glu Pro			
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Val Gly Ser Gln	Glu His Ala Ser Thr	Ala Ser Leu Thr Leu Ile			
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 ttgagntttt tgntaaaaca tggacatgnt tcagtgtgc tontgagaga 200
 gtagcagggtt accacttttg gcaggcccca gccctgcagc aaggaggaag 250
 aggactcaaa agtttggcct ttcactgagc ctccacagca gtgggggaga 300
 agcaagggtt gggcccagtg tcccctttcc ccagtgcac ctcagccttg 350
 gcagccctga taactggtnt ntggctgcaa nttaatgctn tgatatggct 400
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 <210> 78
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